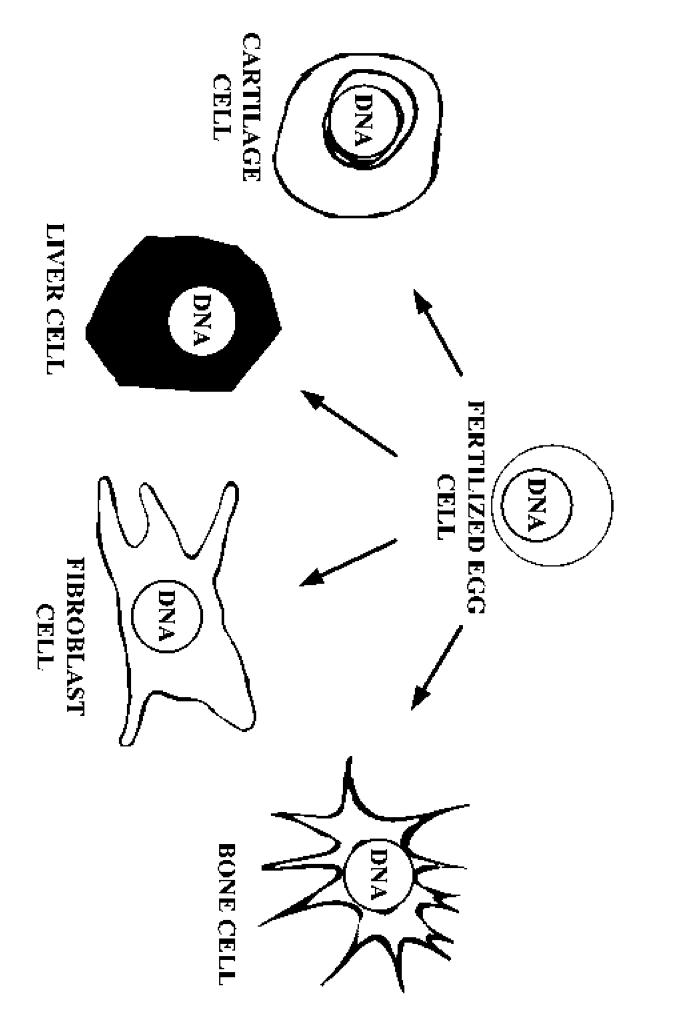
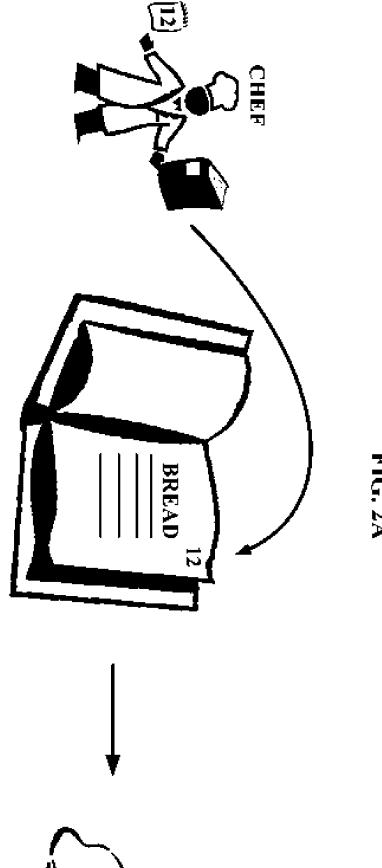
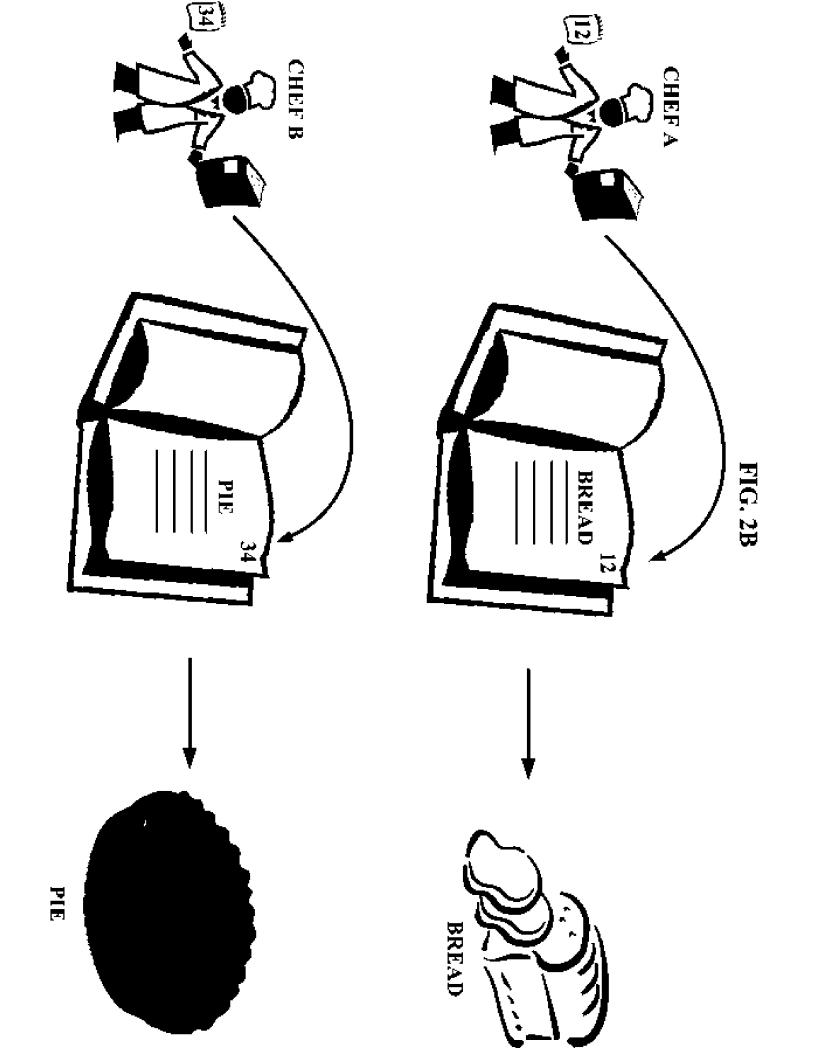
FIG. 1

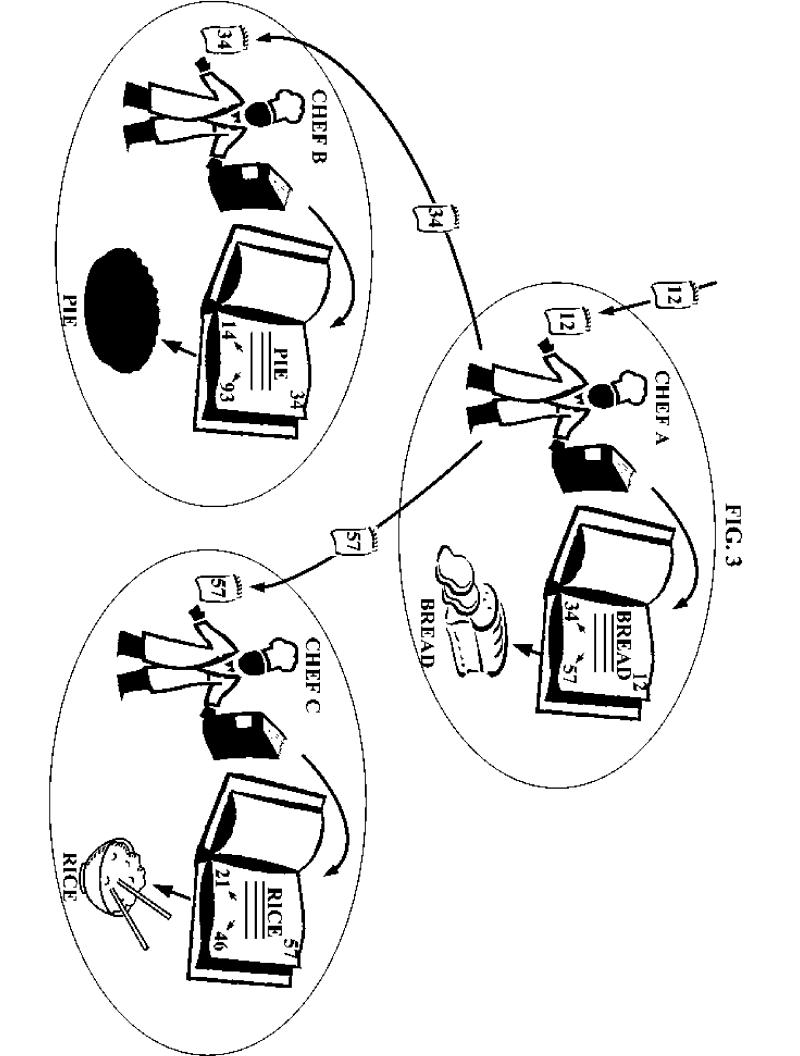




BREAD

FIG. 2A

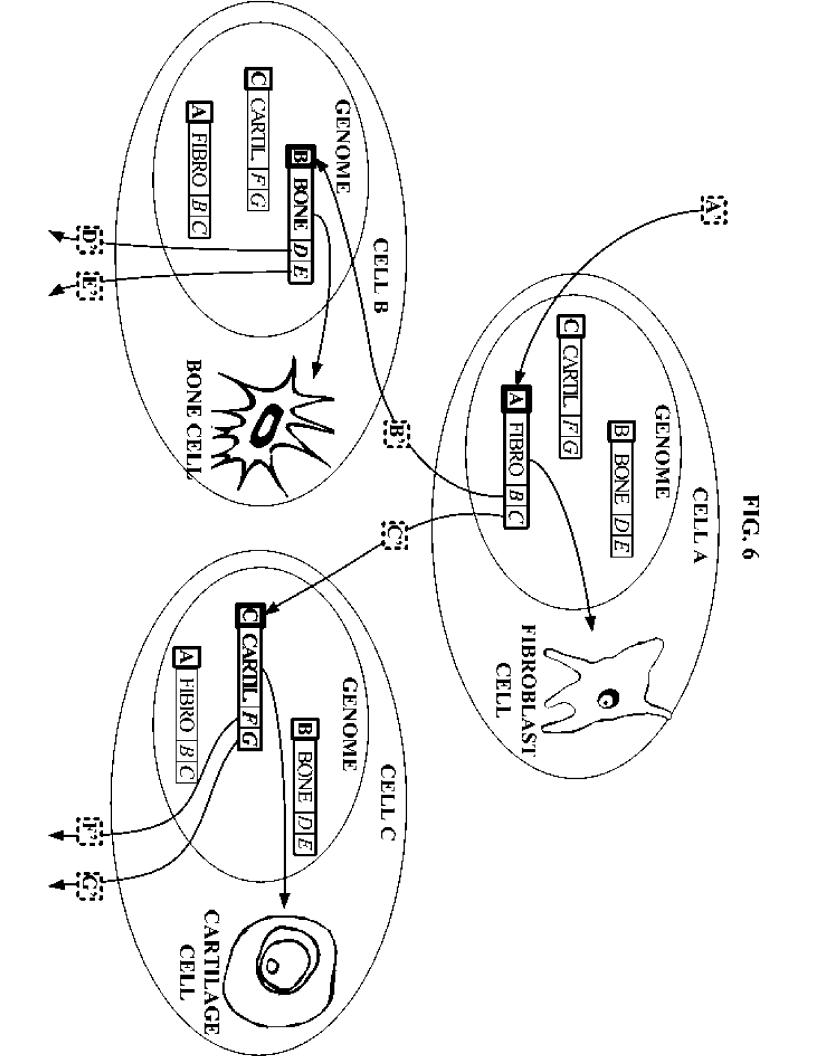


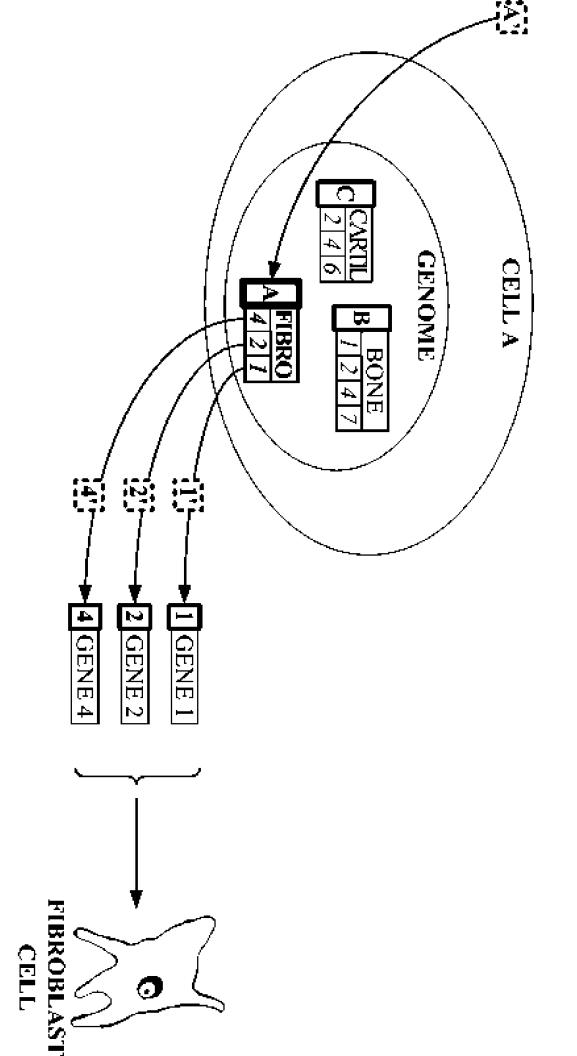


**FIG. 4** 

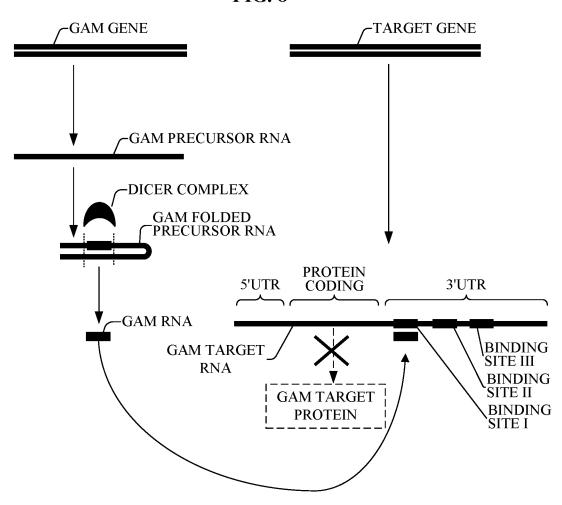
FIG. 5A

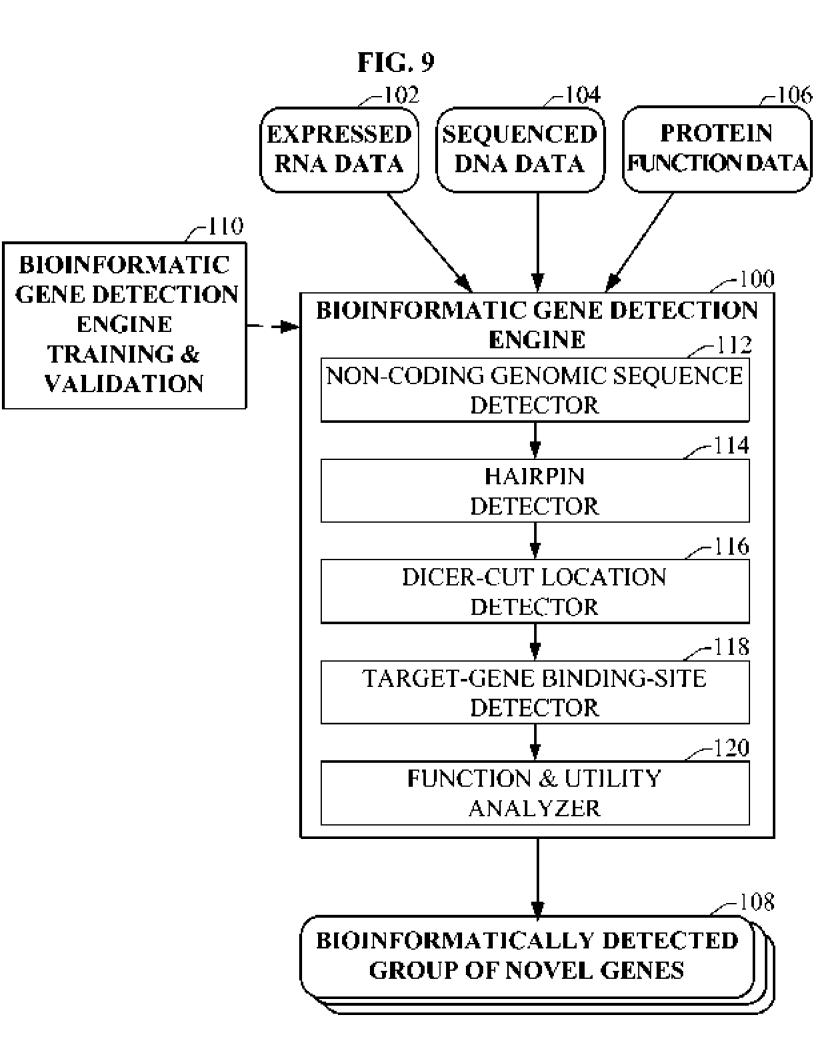
FIG. 5B





**FIG. 8** 





## **FIG. 10**

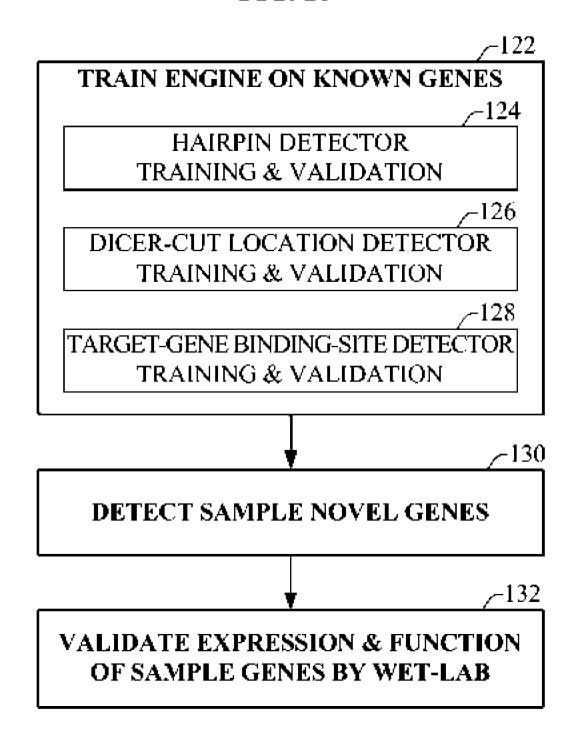


FIG. HA

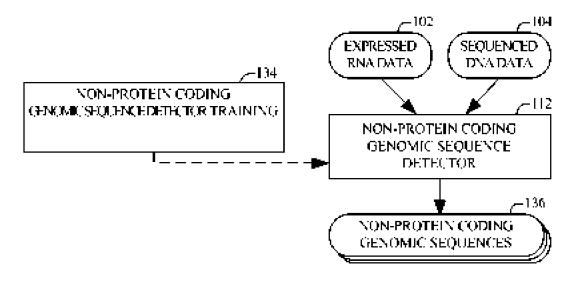
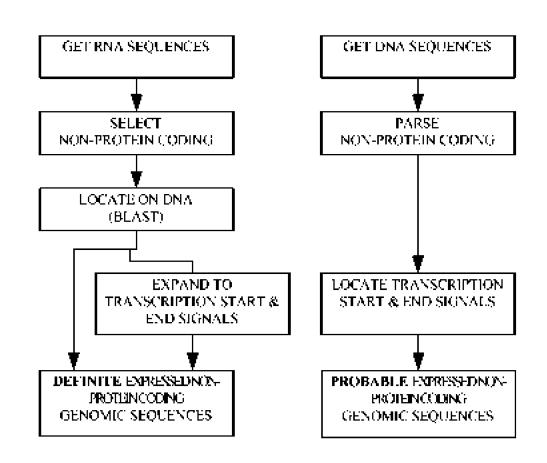
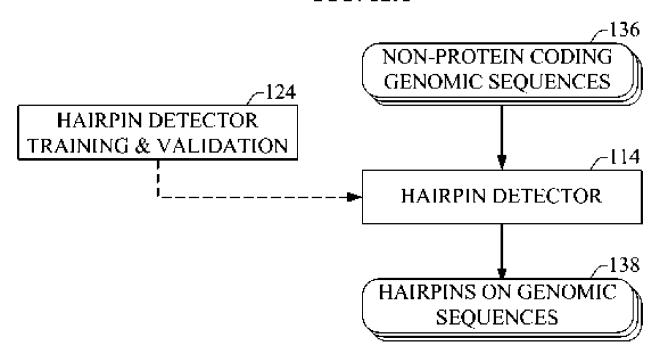


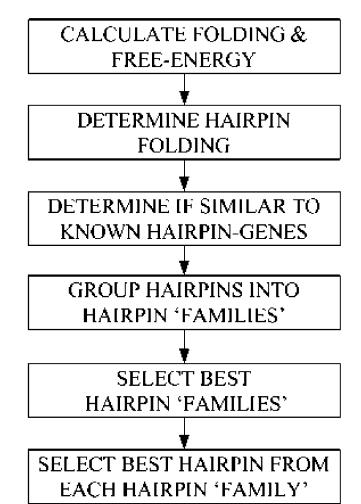
FIG. IIB

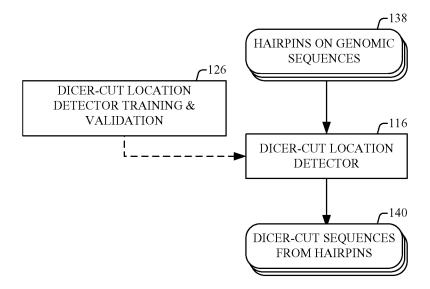


**FIG. 12A** 

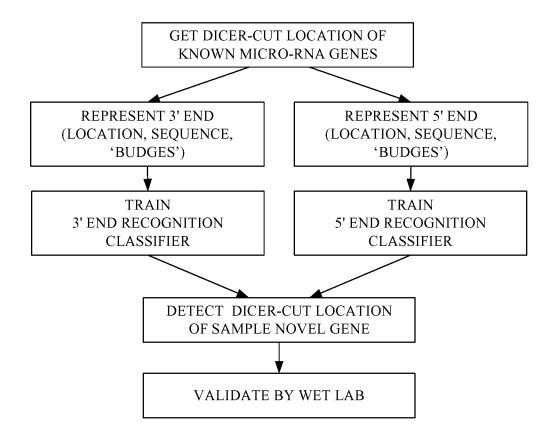


**FIG. 12B** 

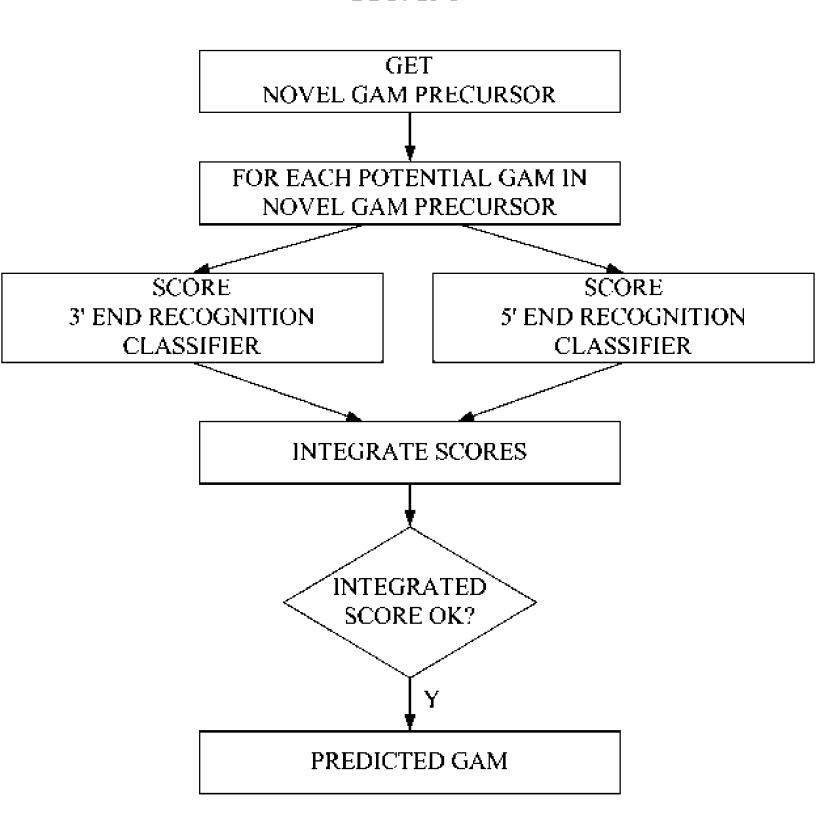




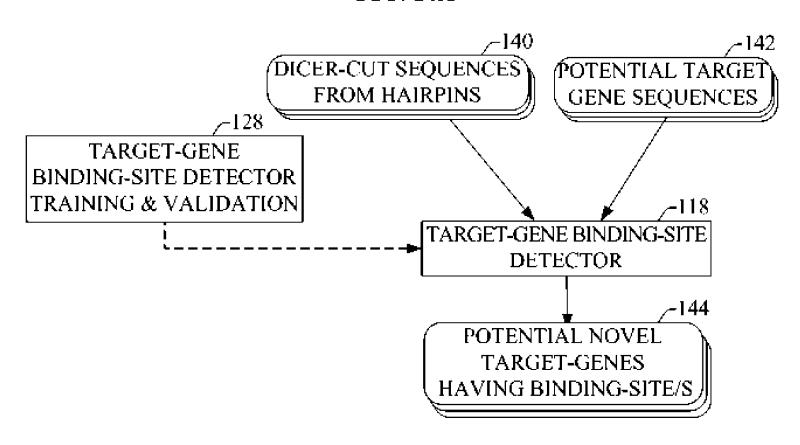
**FIG. 13B** 



**FIG. 13C** 



**FIG. 14A** 



**FIG. 14B** 

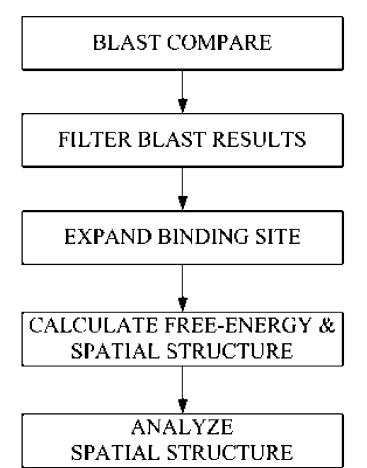
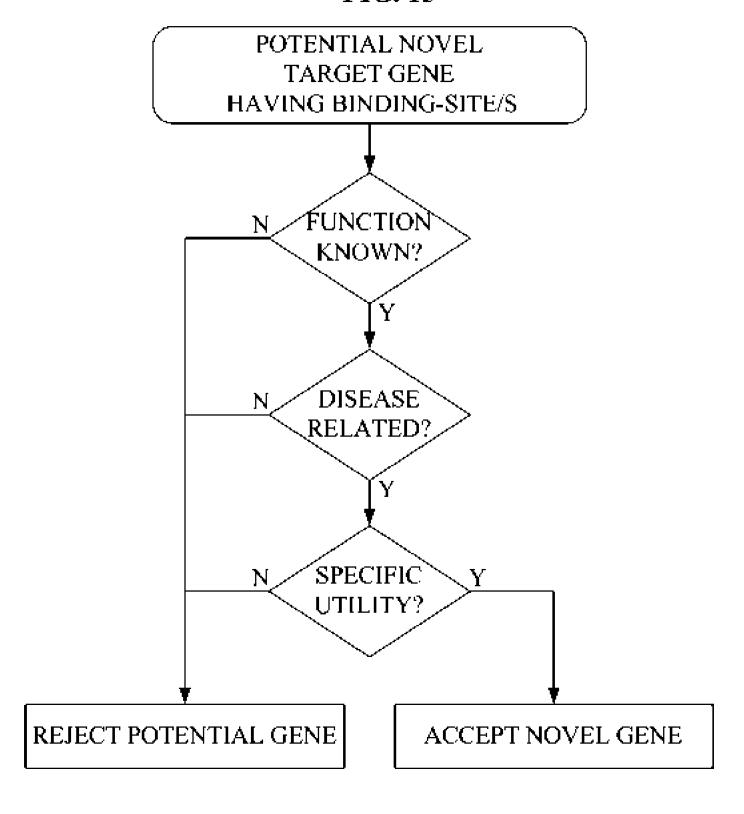


FIG. 15



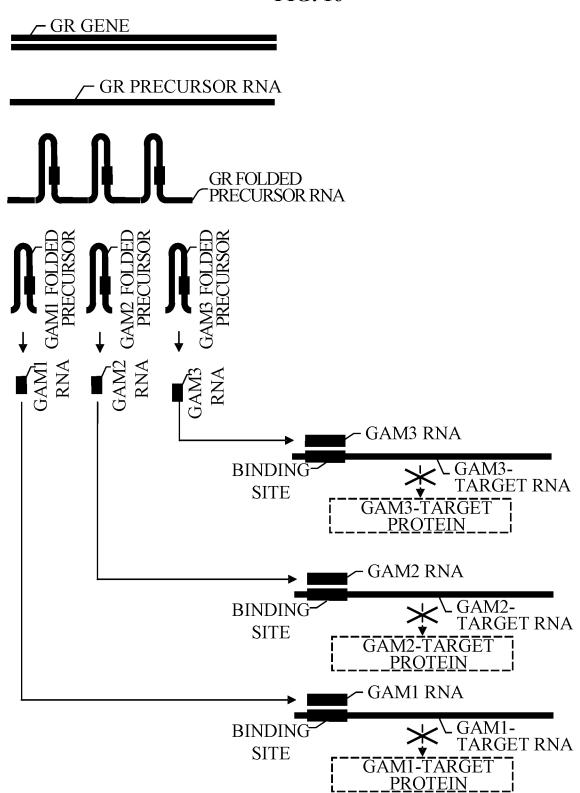


FIG. 18

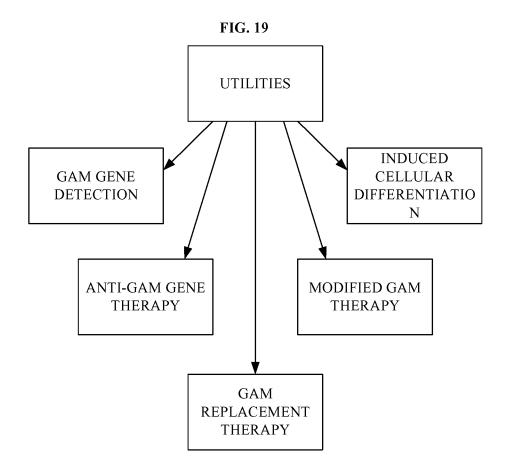
FIND GAM GENES

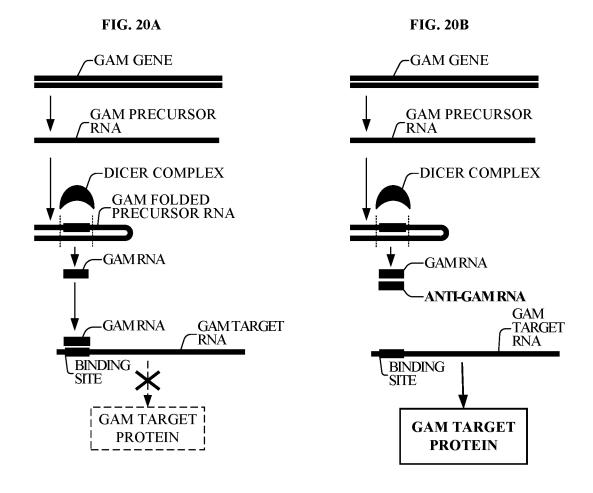
FIND GR GENES

(I.E. GAM GENE CLUSTERS)

FIND HIERARCHY OF
GR GENES

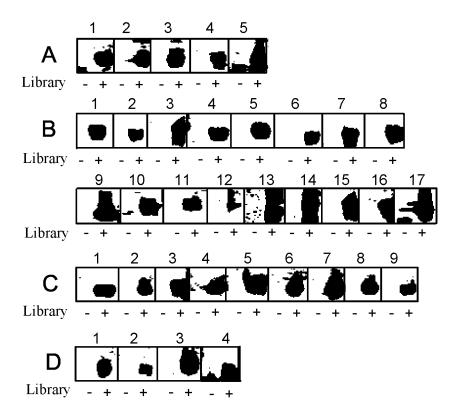
DEDUCE FUNCTION OF
'HIGH' GR AND GAM GENES

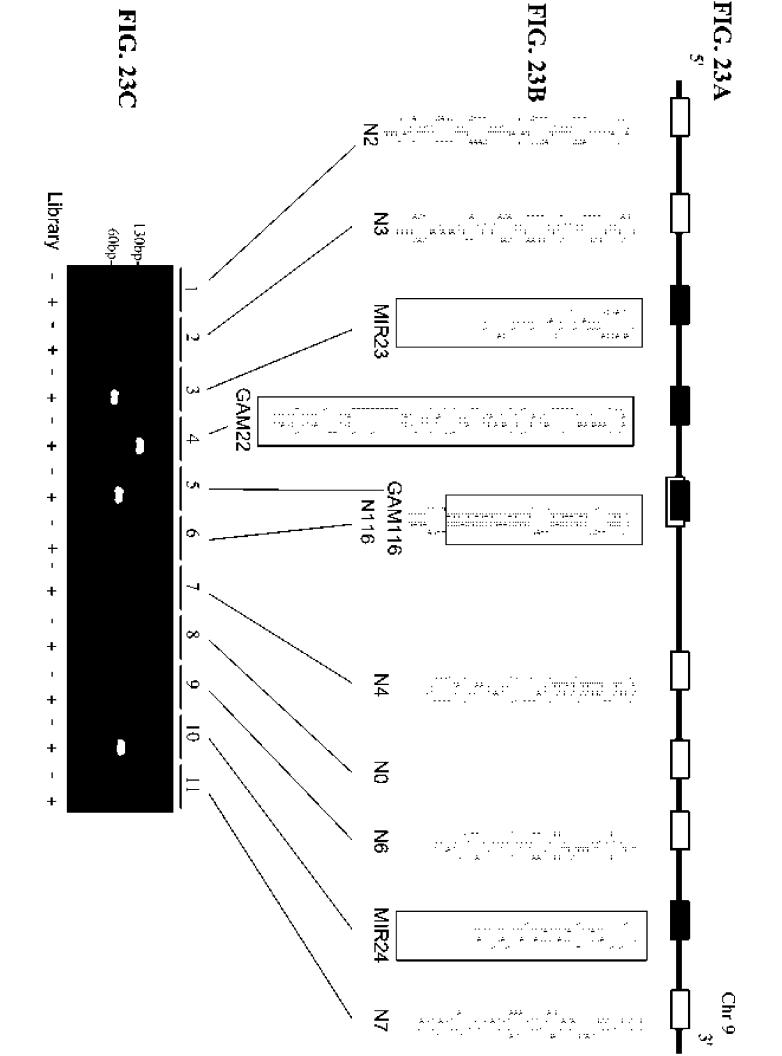




## FIG. 21

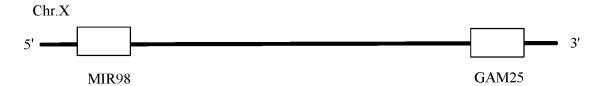
Detection Accuracy Group	Number of Published MIRs	La	Lab Validation	on	Novel GAMs	Hairpins in RNA Database	in RNA base
		Sent	Positive	%		Number	Expected
				success			
A	82	10	5	50%	562	564	282
В	146	91	32	35%	526	813	285
С	38	30	9	30%	625	2080	624
D	78	16	4	25%	3034	11512	2878
Overall	344	147	50	34%	4747	14969	4069



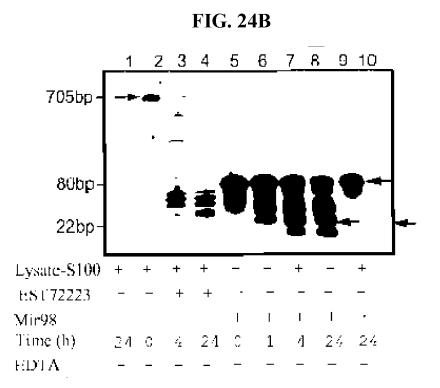


**FIG. 24A** 

## EST72223 (705 nt.)



EST72223 sequence:



## FIG. 24D

